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<210> 20 <211> 418 <212> PRT <213> Homo sapiens

<400> 20

Met Leu Pro Trp Thr Ala Leu Gly Leu Ala Leu Ser Leu Arg Leu Ala 1 10 15 Leu Ala Arg Ser Gly Ala Glu Arg Gly Pro Pro Ala Ser Ala Pro Arg 25 Gly Asp Leu Met Phe Leu Leu Asp Ser Ser Ala Ser Val Ser His Tyr 40 45 Glu Phe Ser Arg Val Arg Glu Phe Val Gly Gln Leu Val Ala Pro Leu 60 Pro Leu Gly Thr Gly Ala Leu Arg Ala Ser Leu Val His Val Gly Ser 70 75 Arg Pro Tyr Thr Glu Phe Pro Phe Gly Gln His Ser Ser Gly Glu Ala 85 90 Ala Gln Asp Ala Val Arg Ala Ser Ala Gln Arg Met Gly Asp Thr His 100 105 110 Thr Gly Leu Ala Leu Val Tyr Ala Lys Glu Gln Leu Phe Ala Glu Ala 115 120

Ser Gly Ala Arg Pro Gly Val Pro Lys Val Leu Val Trp Val Thr Asp 130 135 140 Gly Gly Ser Ser Asp Pro Val Gly Pro Pro Met Gln Glu Leu Lys Asp 150 155 160 Leu Gly Val Thr Val Phe Ile Val Ser Thr Gly Arg Gly Asn Phe Leu 165 170 Glu Leu Ser Ala Ala Ala Ser Ala Pro Ala Glu Lys His Leu His Phe 180 185 Val Asp Val Asp Asp Leu His Ile Ile Val Gln Glu Leu Arg Gly Ser 200 205 Ile Leu Asp Ala Met Arg Pro Gln Gln Leu His Ala Thr Glu Ile Thr 215 220 Ser Ser Gly Phe Arg Leu Ala Trp Pro Pro Leu Leu Thr Ala Asp Ser 230 235 Gly Tyr Tyr Val Leu Glu Leu Val Pro Ser Ala Gln Pro Gly Ala Ala 245 250 Arg Arg Gln Gln Leu Pro Gly Asn Ala Thr Asp Trp Ile Trp Ala Gly 265 270 Leu Asp Pro Asp Thr Asp Tyr Asp Val Ala Leu Val Pro Glu Ser Asn 275 280 Val Arg Leu Leu Arg Pro Gln Ile Leu Arg Val Arg Thr Arg Pro Glu 290 295 300 Glu Ala Gly Pro Glu Arg Ile Val Ile Ser His Ala Arg Pro Arg Ser 310 Leu Arg Val Ser Trp Ala Pro Ala Leu Gly Ser Ala Ala Ala Leu Gly 325 330 Tyr His Val Gln Phe Gly Pro Leu Arg Gly Glu Ala Gln Arg Val 345 Glu Val Pro Ala Gly Arg Asn Cys Thr Thr Leu Gln Gly Leu Ala Pro 360 Gly Thr Ala Tyr Leu Val Thr Val Thr Ala Ala Phe Arg Ser Gly Arg Glu Ser Ala Leu Ser Ala Lys Ala Cys Thr Pro Asp Gly Pro Arg Pro 390 395 Arg Pro Arg Pro Val Pro Arg Ala Pro Thr Pro Gly Thr Ala Ser Arg Glu Pro

<210> 21 <211> 415 <212> PRT <213> Mus musculus

<400> 21

 Met
 Leu
 Phe
 Trp
 Thr
 Ala
 Phe
 Ser
 Met
 Ala
 Leu
 Ser
 Leu
 Arg
 Leu
 Ala
 Leu
 Ala
 Leu
 Arg
 Cly
 Ser
 Thr
 Ala
 Ser
 Asp
 Pro
 Gln
 Arg
 Gln
 Ser
 Arg
 Ala
 Ser
 Ser
 Ala
 Ser
 Val
 Ser
 Ala
 Ser
 Val
 Ala
 Ser
 Ala
 Thr
 Ala
 Ala
 Thr
 Met
 Ala
 Ala
 Thr
 Met
 Ala
 Ala</th

```
100
                                105
Thr Gly Leu Ala Leu Ala Tyr Ala Lys Glu Gln Leu Phe Ala Glu Glu
        115
                            120
Ala Gly Ala Arg Pro Gly Val Pro Lys Val Leu Val Trp Val Thr Asp
                        135
Gly Gly Ser Ser Asp Pro Val Gly Pro Pro Met Gln Glu Leu Lys Asp
145
                    150
                                         155
Leu Gly Val Thr Ile Phe Ile Val Ser Thr Gly Arg Gly Asn Leu Leu
                165
                                     170
Glu Leu Leu Ala Ala Ala Ser Ala Pro Ala Glu Lys His Leu His Phe
                                 185
Val Asp Val Asp Asp Leu Pro Ile Ile Ala Arg Glu Leu Arg Gly Ser
        195
                             200
Ile Thr Asp Ala Met Gln Pro Gln Gln Leu His Ala Ser Glu Val Leu
                        215
Ser Ser Gly Phe Arg Leu Ser Trp Pro Pro Leu Leu Thr Ala Asp Ser
                    230
Gly Tyr Tyr Val Leu Glu Leu Val Pro Ser Gly Lys Leu Ala Thr Thr
                                     250
                245
Arg Arg Gln Gln Leu Pro Gly Asn Ala Thr Ser Trp Thr Trp Thr Asp
                                265
Leu Asp Pro Asp Thr Asp Tyr Glu Val Ser Leu Leu Pro Glu Ser Asn
                            280
Val His Leu Leu Arg Pro Gln His Val Arg Val Arg Thr Leu Gln Glu
                        295
                                             300
Glu Ala Gly Pro Glu Arg Ile Val Ile Ser His Ala Arg Pro Arg Ser
                    310
                                         315
Leu Arg Val Ser Trp Ala Pro Ala Leu Gly Pro Asp Ser Ala Leu Gly
                                     330
Tyr His Val Gln Leu Gly Pro Leu Gln Gly Gly Ser Leu Glu Arg Val
                                345
Glu Val Pro Ala Gly Gln Asn Ser Thr Thr Val Gln Gly Leu Thr Pro
                            360
                                                 365
Cys Thr Thr Tyr Leu Val Thr Val Thr Ala Ala Phe Arg Ser Gly Arg
                        375
Gln Arg Ala Leu Ser Ala Lys Ala Cys Thr Ala Ser Gly Ala Arg Thr
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                                         395
Arg Ala Pro Gln Ser Met Arg Pro Glu Ala Gly Pro Arg Glu Pro
```

```
<210> 22
<211> 182
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<220>

<223> VA domain from collagen XIV

<400> 22

Ile Ala Asp Ile Val Ile Leu Val Asp Gly Ser Trp Ser Ile Gly Arg 10 Phe Asn Phe Arg Leu Val Arg Leu Phe Leu Glu Asn Leu Val Ser Ala Phe Asn Val Gly Ser Glu Lys Thr Arg Val Gly Leu Ala Gln Tyr Ser Gly Asp Pro Arg Ile Glu Trp His Leu Asn Ala Tyr Gly Thr Lys Asp Ala Val Leu Asp Ala Val Arg Asn Leu Pro Tyr Lys Gly Gly Asn Thr

<212> PRT

<213> Artifcial sequence

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70
                                         75
Leu Thr Gly Leu Ala Leu Thr Tyr Ile Leu Glu Asn Ser Phe Lys Pro
                                     90
                                                          95
Glu Ala Gly Ala Arg Pro Gly Val Ser Lys Ile Gly Ile Leu Ile Thr
            100
                                 105
Asp Gly Lys Ser Gln Asp Asp Val Ile Pro Pro Ala Lys Asn Leu Arg
        115
                             120
                                                 125
Asp Ala Gly Ile Glu Leu Phe Ala Ile Gly Val Lys Asn Ala Asp Ile
    1.30
                        135
                                             140
Asn Glu Leu Lys Glu Ile Ala Ser Glu Pro Asp Ser Thr His Val Tyr
                    150
                                         155
Asn Val Ala Asp Phe Asn Phe Met Asn Ser Ile Val Glu Gly Leu Thr
                165
                                     170
Arg Thr Val Cys Ser Arg
            180
<210> 23
<211> 183
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from collagen VII
<400> 23
Ala Ala Asp Ile Val Phe Leu Leu Asp Gly Ser Ser Ser Ile Gly Arg
1
                                     10
Ser Asn Phe Arg Glu Val Arg Ser Phe Leu Glu Gly Leu Val Leu Pro
            20
Phe Ser Gly Ala Ala Ser Ala Gln Gly Val Arg Phe Ala Thr Val Gln
Tyr Ser Asp Asp Pro Arg Thr Glu Phe Gly Leu Asp Ala Leu Gly Ser
                        55
Gly Gly Asp Val Ile Arg Ala Ile Arg Glu Leu Ser Tyr Lys Gly Gly
                    70
Asn Thr Arg Thr Gly Ala Ala Ile Leu His Val Ala Asp His Val Phe
                85
                                     90
Leu Pro Gln Leu Ala Arg Pro Gly Val Pro Lys Val Cys Ile Leu Ile
                                 105
                                                     110
Thr Asp Gly Lys Ser Gln Asp Leu Val Asp Thr Ala Ala Gln Arg Leu
        115
                            120
Lys Gly Gln Gly Val Lys Leu Phe Ala Val Gly Ile Lys Asn Ala Asp
    130
                                             140
Pro Glu Glu Leu Lys Arg Val Ala Ser Gln Pro Thr Ser Asp Phe Phe
                    150
                                         155
Phe Phe Val Asn Asp Phe Ser Ile Leu Arg Thr Leu Leu Pro Leu Val
                165
                                     170
Ser Arg Arg Val Cys Thr Thr
            180
<210> 24
<211> 182
<212> PRT
<213> Artificial Sequence
```

<223> VA domain from collagen XII

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<400> 24
Lys Ala Asp Ile Val Phe Leu Thr Asp Ala Ser Trp Ser Ile Gly Asp
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Asp Asn Phe Asn Lys Val Val Lys Phe Ile Phe Asn Thr Val Gly Ala
            2.0
Phe Asp Glu Val Asn Pro Ala Gly Ile Gln Val Ser Phe Val Gln Tyr
                             40
Ser Asp Glu Val Lys Ser Glu Phe Lys Leu Asn Thr Tyr Asn Asp Lys
    50
                        55
Ala Leu Ala Leu Gly Ala Leu Gln Asn Ile Arg Tyr Arg Gly Gly Asn
                    70
                                         75
Thr Arg Thr Gly Lys Ala Leu Thr Phe Ile Lys Glu Lys Val Leu Thr
                85
                                     90
Trp Glu Ser Gly Met Arg Lys Asn Val Arg Val Leu Gly Val Val Thr
            100
                                 105
Asp Gly Arg Ser Gln Asp Glu Val Lys Lys Ala Ala Phe Val Ile Gln
        115
                             120
Gln Ser Gly Phe Ser Val Phe Val Val Gly Val Ala Asp Val Asp Tyr
    130
                        135
Asn Glu Leu Ala Asn Ile Ala Ser Lys Pro Ser Glu Arg His Val Phe
                    150
Ile Val Asp Asp Phe Glu Ser Phe Glu Lys Ile Glu Asp Asn Leu Ile
                165
                                     170
Thr Phe Val Cys Glu Thr
            180
<210> 25
<211> 185
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from collagen VI
<400> 25
Ala Ala Asp Ile Val Phe Leu Val Asp Ser Ser Trp Ser Ala Gly Lys
1
Asp Arg Phe Leu Leu Val Gln Glu Phe Leu Ser Asp Val Val Glu Ser
            20
Leu Ala Val Gly Asp Asn Asp Phe His Phe Ala Leu Val Arg Leu Asn
Gly Asn Pro His Thr Glu Phe Leu Leu Asn Thr Tyr His Ser Lys Gln
Glu Val Leu Ser His Ile Ala Asn Met Ser Tyr Ile Gly Gly Ser Asn
                                         75
Gln Thr Gly Lys Gly Leu Glu Tyr Val Ile His Ser His Leu Thr Glu
                                    90
Ala Ser Gly Ser Arg Ala Ala Asp Gly Val Pro Gln Val Ile Val Val
                                105
Leu Thr Asp Gly Gln Ser Glu Asp Gly Phe Ala Leu Pro Ser Ala Glu
                            120
        115
```

Leu Lys Ser Ala Asp Val Asn Val Phe Ala Val Gly Val Glu Gly Ala

Asp Glu Arg Ala Leu Gly Glu Val Ala Ser Glu Pro Leu Leu Ser Met

His Val Phe Asn Leu Glu Asn Val Thr Ser Leu His Gly Leu Val Gly

155

170

135

150

130

```
180
<210> 26
<211> 185
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from matrilin-2
<400> 26
Arg Ala Asp Leu Val Phe Ile Ile Asp Ser Ser Arg Ser Val Asn Thr
1
                                     10
Tyr Asp Tyr Ala Lys Val Lys Glu Phe Ile Leu Asp Ile Leu Gln Phe
            20
                                25
Leu Asp Ile Gly Pro Asp Val Thr Arg Val Gly Leu Leu Gln Tyr Gly
        35
                             40
Ser Thr Val Lys Asn Glu Phe Ser Leu Lys Thr Phe Lys Arg Lys Ser
                        55
Glu Val Glu Arg Ala Val Lys Arg Met Arg His Leu Ser Thr Gly Thr
                                         75
Met Thr Gly Leu Ala Ile Gln Tyr Ala Leu Asn Ile Ala Phe Ser Glu
                                     90
Ala Glu Gly Ala Arg Pro Leu Arg Glu Asn Val Pro Arg Ile Ile Met
            100
                                105
Ile Val Thr Asp Gly Arg Pro Gln Asp Ser Val Ala Glu Val Ala Ala
                            120
Lys Ala Arg Asn Thr Gly Ile Leu Ile Phe Ala Ile Gly Val Gly Gln
                        135
Val Asp Leu Asn Thr Leu Lys Ala Ile Gly Ser Glu Pro His Lys Asp
                    150
                                         155
His Val Phe Leu Val Ala Asn Phe Ser Gln Ile Glu Ser Leu Thr Ser
                165
                                    170
Val Phe Gln Asn Lys Leu Cys Thr Val
            180
<210> 27
<211> 184
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from matrilin-4
<400> 27
Pro Leu Asp Leu Val Phe Met Ile Asp Ser Ser Arg Ser Val Arg Pro
                                    10
Phe Glu Phe Glu Thr Met Arg Gln Phe Leu Val Gly Leu Leu Arg Ser
Leu Asp Val Gly Leu Asn Ala Thr Arg Val Gly Val Ile Gln Tyr Ser
Ser Gln Val Gln Ser Val Phe Pro Leu Gly Ala Phe Ser Arg Arg Glu
                        55
Asp Met Glu Arg Ala Ile Arg Ala Val Val Pro Leu Ala Gln Gly Thr
Met Thr Gly Leu Ala Ile Gln Tyr Ala Met Asn Val Ala Phe Ser Glu
```

Asn Leu Val Ser Cys Ile His Ser Ser

```
90
                85
Ala Glu Gly Ala Arg Pro Ser Glu Glu Arg Val Pro Arg Val Leu Val
            100
                                 105
Ile Val Thr Asp Gly Arg Pro Gln Asp Arg Val Ala Glu Val Ala Ala
        115
                             120
Gln Ala Arg Ala Arg Gly Ile Glu Ile Tyr Ala Val Gly Val Gln Arg
                        135
                                             140
Ala Asp Val Gly Ser Leu Arg Thr Met Ala Ser Pro Pro Leu Asp Gln
                    150
                                         155
His Val Phe Leu Val Glu Ser Phe Asp Ile Gln Glu Phe Gly Leu Gln
                165
                                     170
Phe Gln Gly Arg Leu Cys Gly Lys
            180
<210> 28
<211> 185
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from matrilin-3
<400> 28
Pro Leu Asp Leu Val Phe Ile Ile Asp Ser Ser Arg Ser Val Arg Pro
1
                                     10
Leu Glu Phe Thr Lys Val Lys Thr Phe Val Ser Arg Ile Ile Asp Thr
Leu Asp Ile Gly Ala Thr Asp Thr Arg Val Ala Val Val Asn Tyr Ala
                            40
Ser Thr Val Lys Ile Glu Phe Gln Leu Asn Thr Tyr Ser Asp Lys Gln
                        55
Ala Leu Lys Gln Ala Val Ala Arg Ile Thr Pro Leu Ser Thr Gly Thr
                    70
Met Ser Gly Leu Ala Ile Gln Thr Ala Met Glu Glu Ala Phe Thr Val
                                    90
Glu Ala Gly Ala Arg Gly Pro Met Ser Asn Ile Pro Lys Val Ala Ile
                                105
Ile Val Thr Asp Gly Arg Pro Gln Asp Gln Val Asn Glu Val Ala Ala
        115
                            120
Arg Ala Arg Ala Ser Gly Ile Glu Leu Tyr Ala Val Gly Val Asp Arg
                        135
Ala Asp Met Glu Ser Leu Lys Met Met Ala Ser Lys Pro Leu Glu Glu
                    150
                                         155
His Val Phe Tyr Val Glu Thr Tyr Gly Val Ile Glu Lys Leu Ser Ala
                165
                                    170
Arg Phe Gln Glu Thr Pro Cys Ala Leu
            180
<210> 29
<211> 185
<212> PRT
<213> Artificial Sequence
<223> VA domain from matrilin-1
<400> 29
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Pro Thr Asp Leu Val Phe Val Val Asp Ser Ser Arg Ser Val Arg Pro 10 Val Glu Phe Glu Lys Val Lys Val Phe Leu Ser Gln Val Ile Glu Ser Leu Asp Val Gly Pro Asn Ala Thr Arg Val Gly Leu Val Asn Tyr Ala 40 Ser Thr Val Lys Pro Glu Phe Pro Leu Arg Ala His Gly Ser Lys Ala Ser Leu Leu Gln Ala Val Arg Arg Ile Gln Pro Leu Ser Thr Gly Thr Met Thr Gly Leu Ala Leu Gln Phe Ala Ile Thr Lys Ala Leu Ser Asp 90 Ala Glu Gly Gly Arg Ala Arg Ser Pro Asp Ile Ser Lys Val Val Ile 100 105 Val Val Thr Asp Gly Arg Pro Gln Asp Ser Val Arg Asp Val Ser Glu 120 Arg Ala Arg Ala Ser Gly Ile Glu Leu Phe Ala Ile Gly Leu Gly Arg 135 Val Asp Lys Ala Thr Leu Arg Gln Ile Ala Ser Glu Pro Gln Asp Glu His Val Asp Tyr Val Glu Ser Tyr Asn Val Ile Glu Lys Leu Ala Lys 1.65 Lys Phe Gln Glu Ala Phe Cys Val Val 180

<210> 30 <211> 193 <212> PRT

<213> Artificial Sequence

<220>

<223> VA domain from VLA

<400> 30 Gln Leu Asp Ile Val Ile Val Leu Asp Gly Ser Asn Ser Ile Tyr Pro 10 Trp Asp Ser Val Thr Ala Phe Leu Asn Asp Leu Leu Lys Arg Met Asp Ile Gly Pro Lys Gln Thr Gln Val Gly Ile Val Gln Tyr Gly Glu Asn Val Thr His Glu Phe Asn Leu Asn Lys Tyr Ser Ser Thr Glu Glu Val Leu Val Ala Ala Lys Lys Ile Val Gln Arg Gly Gly Arg Gln Thr Met Thr Ala Leu Gly Thr Asp Thr Ala Arg Lys Glu Ala Phe Thr Glu Ala 90 Arg Gly Ala Arg Arg Gly Val Lys Lys Val Met Val Ile Val Thr Asp 100 Gly Glu Ser His Asp Asn His Arg Leu Lys Lys Val Ile Gln Asp Cys 120 Glu Asp Glu Asn Ile Gln Arg Phe Ser Ile Ala Ile Leu Gly Ser Tyr 135 Asn Arg Gly Asn Leu Ser Thr Glu Lys Phe Val Glu Glu Ile Lys Ser 150 Ile Ala Ser Glu Pro Thr Glu Lys His Phe Phe Asn Val Ser Asp Glu 165 170 Leu Ala Leu Val Thr Ile Val Lys Thr Leu Gly Glu Arg Ile Phe Ala

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<210> 31
<211> 181
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from WARP
<400> 31
Gln Gly Asp Leu Leu Phe Leu Leu Asp Ser Ser Ala Ser Val Ser His
Tyr Glu Phe Ser Arg Val Arg Glu Phe Val Gly Gln Leu Val Ala Thr
            20
Met Ser Phe Gly Pro Gly Ala Leu Arg Ala Ser Leu Val His Val Gly
Ser Gln Pro His Thr Glu Phe Thr Phe Asp Gln Tyr Ser Ser Gly Gln
Ala Ile Arg Asp Ala Ile Arg Val Ala Pro Gln Arg Met Gly Asp Thr
                                         75
Asn Thr Gly Leu Ala Leu Ala Tyr Ala Lys Glu Gln Leu Phe Ala Glu
                                     90
Glu Ala Gly Ala Arg Pro Gly Val Pro Lys Val Leu Val Trp Val Thr
                                105
                                                     110
Asp Gly Gly Ser Ser Asp Pro Val Gly Pro Pro Met Gln Glu Leu Lys
                            120
Asp Leu Gly Val Thr Ile Phe Ile Val Ser Thr Gly Arg Gly Asn Leu
                        135
                                             140
Leu Glu Leu Leu Ala Ala Ser Ala Pro Ala Glu Lys His Leu His
                    150
                                         155
Phe Val Asp Val Asp Asp Leu Pro Ile Ile Ala Arg Glu Leu Arg Gly
                                    170
Ser Ile Thr Asp Ala
            180
<210> 32
<211> 184
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from cochlin
<400> 32
Lys Ala Asp Ile Ala Phe Leu Ile Asp Gly Ser Tyr Asn Ile Gly Gln
1
                                    10
Arg Arg Phe Asn Leu Gln Lys Asn Phe Val Gly Lys Val Ala Val Met
                                25
Leu Gly Ile Gly Thr Glu Gly Pro His Val Gly Val Val Gln Ala Ser
Glu His Pro Lys Ile Glu Phe Tyr Leu Lys Asn Phe Thr Ala Ala Lys
Glu Val Leu Phe Ala Ile Lys Glu Leu Gly Phe Arg Gly Gly Asn Ser
Asn Thr Gly Lys Ala Leu Lys His Ala Ala Gln Lys Phe Phe Ser Met
```

```
90
Glu Asn Gly Ala Arg Lys Gly Ile Pro Lys Ile Ile Val Val Phe Leu
            100
                                105
Asp Gly Trp Pro Ser Asp Asp Leu Glu Glu Ala Gly Ile Val Ala Arg
        115
                            120
                                                 125
Glu Phe Gly Val Asn Val Phe Ile Val Ser Ser Val Ala Lys Pro Thr
                        135
                                             140
Thr Glu Glu Leu Gly Met Val Gln Asp Ile Gly Phe Ile Asp Lys Ala
                    150
                                         155
Val Cys Arg Asn Asn Gly Phe Phe Ser Tyr Gln Met Pro Ser Trp Phe
                                     170
                165
Gly Thr Thr Lys Tyr Val Lys Pro
            180
<210> 33
<211> 186
<212> PRT
<213> Artificial Sequence
<220>
<223> VA domain from vwf
<400> 33
Leu Leu Asp Leu Val Phe Leu Leu Asp Gly Ser Ser Arg Leu Ser Glu
1
Ala Glu Phe Glu Val Leu Lys Ala Phe Val Val Asp Met Met Glu Arq
Leu Arg Ile Ser Gln Lys Trp Val Arg Val Ala Val Val Glu Tyr His
Asp Gly Ser His Ala Tyr Ile Gly Leu Lys Asp Arg Lys Arg Pro Ser
    50
                        55
Glu Leu Arg Arg Ile Ala Ser Gln Val Lys Tyr Ala Gly Ser Gln Val
                    70
                                         75
Ala Ser Thr Ser Glu Val Leu Lys Tyr Thr Leu Phe Gln Ile Phe Ser
                                    90
Lys Ile Asp Arg Pro Glu Ala Ser Arg Ile Ala Leu Leu Leu Met Ala
            100
                                105
                                                     110
Ser Gln Glu Pro Gln Arg Met Ser Arg Asn Phe Val Arg Tyr Val Gln
                            120
                                                 125
Gly Leu Lys Lys Lys Val Ile Val Ile Pro Val Gly Ile Gly Pro
    130
                        135
His Ala Asn Leu Lys Gln Ile Arg Leu Ile Glu Lys Gln Ala Pro Glu
                    150
Asn Lys Ala Phe Val Leu Ser Ser Val Asp Glu Leu Glu Gln Gln Arg
                165
                                    170
Asp Glu Ile Val Ser Tyr Leu Cys Asp Leu
            180
<210> 34
<211> 85
<212> PRT
<213> Artificial Sequence
<223> F3-3 repeats from collagen XII
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<400> 34

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Pro Arg Asn Leu Lys Val Thr Asp Glu Thr Thr Asp Ser Phe Lys Ile
Thr Trp Thr Gln Ala Pro Gly Arg Val Leu Arg Tyr Arg Ile Ile Tyr
                                25
Arg Pro Val Ala Gly Gly Glu Ser Arg Glu Val Thr Thr Pro Pro Asn
                            40
Gln Arg Arg Thr Leu Glu Asn Leu Ile Pro Asp Thr Lys Tyr Glu
                        55
Val Ser Val Ile Pro Glu Tyr Phe Ser Gly Pro Gly Thr Pro Leu Thr
Gly Asn Ala Ala Thr
<210> 35
<211> 86
<212> PRT
<213> Artificial Sequence
<223> F3-12 repeats from fibronectin
<400> 35
Pro Ser Gln Met Gln Val Thr Asp Val Gln Asp Asn Ser Ile Ser Val
1
                                    10
Arg Trp Leu Pro Ser Thr Ser Pro Val Thr Gly Tyr Arg Val Thr Thr
            20
Thr Pro Lys Asn Gly Leu Gly Pro Ser Lys Thr Lys Thr Ala Ser Pro
        35
                            40
Asp Gln Thr Glu Met Thr Ile Glu Gly Leu Gln Pro Thr Val Glu Tyr
                        55
Val Val Ser Val Tyr Ala Gln Asn Arg Asn Gly Glu Ser Gln Pro Leu
                    70
Val Gln Thr Ala Val Thr
<210> 36
<211> 87
<212> PRT
<213> Artificial Sequence
<220>
<223> F3-2 repeats from WARP
<400> 36
Pro Glu Arg Ile Val Ile Ser His Ala Arg Pro Arg Ser Leu Arg Val
                                    10
Ser Trp Ala Pro Ala Leu Gly Pro Asp Ser Ala Leu Gly Tyr His Val
                                25
                                                    30
Gln Leu Gly Pro Leu Gln Gly Gly Ser Leu Glu Arg Val Glu Val Pro
                            40
Ala Gly Gln Asn Ser Thr Thr Val Gln Gly Leu Thr Pro Cys Thr Thr
                        55
Tyr Leu Val Thr Val Thr Ala Ala Phe Arg Ser Gly Arg Gln Arg Ala
                    70
                                        75
```

Leu Ser Ala Lys Ala Cys Thr

85

```
<210> 37
<211> 88
<212> PRT
<213> Artificial Sequence
<223> F3-3 repeats from beta-4 integrin
<400> 37
Pro Thr Arg Leu Val Phe Ser Ala Leu Gly Pro Thr Ser Leu Arg Val
 1
Ser Trp Gln Glu Pro Arg Cys Glu Arg Pro Leu Gln Gly Tyr Ser Val
            20
Glu Tyr Gln Leu Leu Asn Gly Gly Glu Leu His Arg Leu Asn Ile Pro
                             40
Asn Pro Ala Gln Thr Ser Val Val Val Glu Asp Leu Leu Pro Asn His
                         55
Ser Tyr Val Phe Arg Val Arg Ala Gln Ser Gln Glu Gly Trp Gly Arg
                    70
Glu Arg Glu Gly Val Ile Thr Ile
                85
<210> 38
<211> 85
<212> PRT
<213> Artificial Sequence
<223> F3-5 repeat from collagen XIV
<400> 38
Pro Gln His Leu Glu Val Asp Glu Ala Ser Thr Asp Ser Phe Arg Val
 1
Ser Trp Lys Pro Thr Ser Ser Asp Ile Ala Phe Tyr Arg Leu Ala Trp
Ile Pro Leu Asp Gly Gly Glu Ser Glu Glu Val Val Leu Ser Gly Asp
Ala Asp Ser Tyr Val Ile Glu Gly Leu Leu Pro Asn Thr Glu Tyr Glu
Val Ser Leu Leu Ala Val Phe Asp Asp Glu Thr Glu Ser Glu Val Val
Ala Val Leu Gly Ala
<210> 39
<211> 85
<212> PRT
<213> Artificial Sequence
<223> F3-7 repeat from tenascin-R
<400> 39
Pro Lys Asp Ile Thr Ile Ser Asn Val Thr Lys Asp Ser Val Met Val
Ser Trp Ser Pro Pro Val Ala Ser Phe Asp Tyr Tyr Arg Val Ser Tyr
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```
25
Arg Pro Thr Gln Val Gly Arg Leu Asp Ser Ser Val Val Pro Asn Thr
                            40
                                                45
Val Thr Glu Phe Thr Ile Thr Arg Leu Asn Pro Ala Thr Glu Tyr Glu
                        55
                                            60
Ile Ser Leu Asn Ser Val Arg Gly Arg Glu Glu Ser Glu Arg Ile Cys
                                                             80
Thr Leu Val His Thr
<210> 40
<211> 87
<212> PRT
<213> Artificial Sequence
<220>
<223> F3-1 repeat from WARP
<400> 40
Pro Gln Gln Leu His Ala Ser Glu Val Leu Ser Ser Gly Phe Arg Leu
1
                                    10
Ser Trp Pro Pro Leu Leu Thr Ala Asp Ser Gly Tyr Tyr Val Leu Glu
            20
                                25
Leu Val Pro Ser Gly Lys Leu Ala Thr Thr Arg Arg Gln Gln Leu Pro
                            40
Gly Asn Ala Thr Ser Trp Thr Trp Thr Asp Leu Asp Pro Asp Thr Asp
                        55
                                            60
Tyr Glu Val Ser Leu Leu Pro Glu Ser Asn Val His Leu Leu Arg Pro
65
                    70
                                        75
```

Gln His Val Arg Val Arg Thr

85